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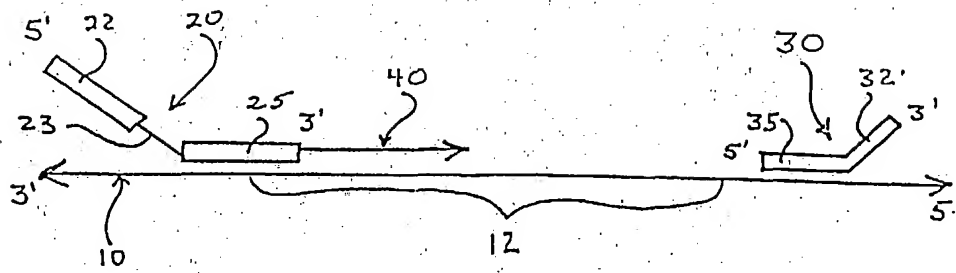


Fig. 1

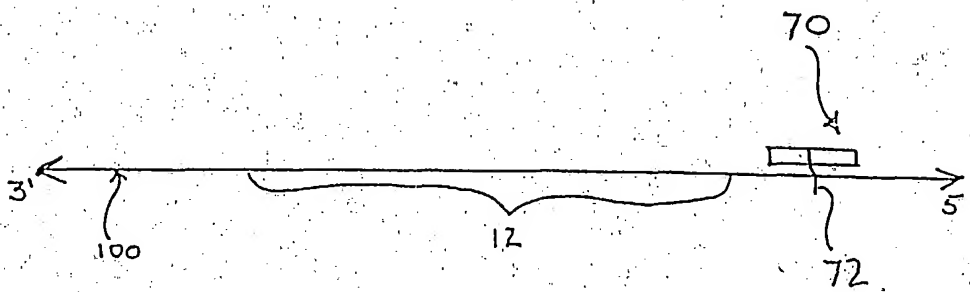


Fig. 2A

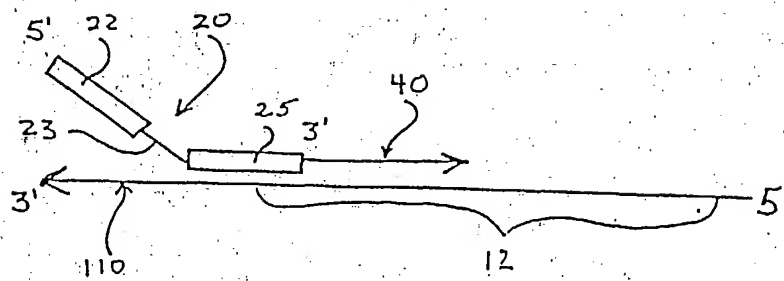


Fig. 2B

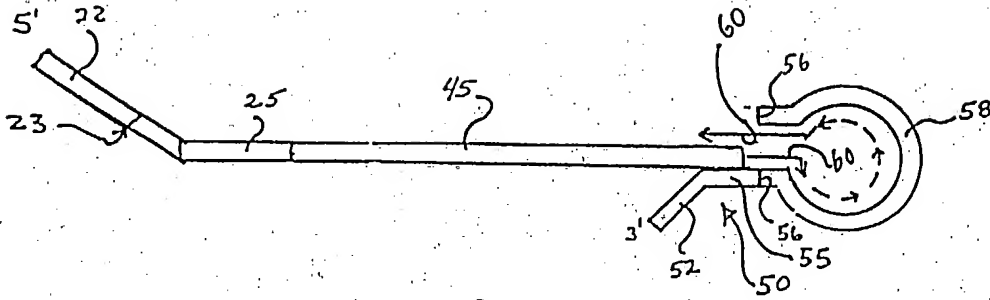


Fig. 3

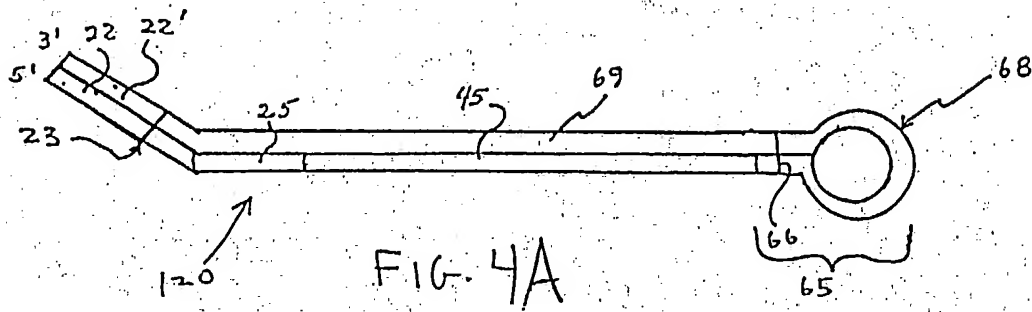


FIG. 4A

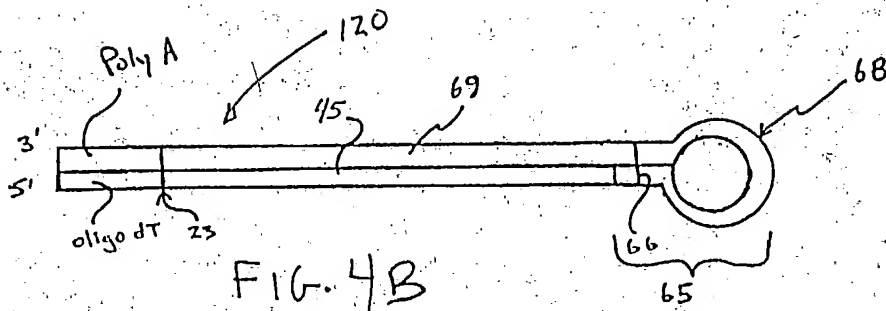


FIG. 4B

clones	SEQ ID NO.	FR1	CDR1	FR2
A7, B1, F8, G2	209	ESDGA VVQPGGSLRLSCAASGF	IFDDFAMH	WLRQVPGKGLQWVGL
C2, E6	210	QPGGSLRLSCAASGF	TLSSSAMS	WVRQAPGKGLEFVAV
A4	211	QPGGSLRLSCAASGF	TLSSSAMS	WVRQAPGKGLEFVAV
F6	212	AWYSRGSPCLSCAASGF	TLSSSAMS	WVRQAPGKGLEFVAV
E9, G7	213	ESDPGLVKPSETPSLTCTVSGG	SISSTMVFWG	WIRQPPGKGLEWIAS
F3, G4	214	PGLVKPSETPSLTCTVSGG	SISNMVFWG	WIRQPPGKGLEWIAS
A12, B5, B8, B9	215	ESDPGLVQPSQTLSTCTVSGG	SLRSDYYWS	WIRQSPGKGLEWIAI
E8	216	PVQPLEF	TFTDHWMH	WVRQAPGKGLVWLAR
F7	217	ESEGLVQPGGSLRLSCAASGF	TFSSYAMT	WVRQAPGKGLEWVST
E11	218	LAGVEVVQPGGSLRLSCAASGF	TFDDYAMH	WLRQIPGKGLQWVSL

Figure 5A

clones	CDR2	FR3
A7, B1, F8, G2	MSWDGVSAYYADSVEG	RFTISRDNKKNALYLQMSLIGVEDTALYYCAK
C2, E6	SSGNGFSTYYGDSVKG	RFTISRDNCKMWVYLQMSLRAEDTAKYHCAK
A4	SSGNGFSTYYGDSVKG	RFTISRDNCKMWVYLQMSLRAEDTAKYHCAK
F6	SSGNGFSTYYGDSVKG	RFTISRDNCKMWVYLQMSLRAEDTAKYHCAK
E9, G7	IYYSGTT-YYNPSLRS	RVTMSVDTSKNQLSLKLSVTAADTAVYYCAR
F3, G4	IYYSGTT-YYNPSLRS	RVTMSVDTSKNQLSLKLSVTAADTAVYYCAR
A12, B5, B8, B9	ISYTGTT-YYNPSLKS	RVTISVDTSRNQFSLRLRSVTAADS AVYFCAS
E8	INRDGSDTTYADSVTG	RFTISRDNKGKNTVSLQMSLSVDDTAVYYCAR
F7	MTGSGGVTTYADVLKG	RFTISRDNCKNTLYLQMSLRAEDTAVYYCAK
E11	LSWDGVSAYYADSVEG	RFTISRDNKKNSLYLQMSLRAEDVALYYCAK

Figure 5B

clones	CDR3	FR4
A7, B1, F8, G2	DMGGGLRFPHF	WGQGTPLVTSA
C2, E6	VRYPGRSHFFFD	WGQGTPLVTSS
A4	VRYPGRSHFFFD	WPGNPGHRL
F6	VRYPGRSHFFFD	WGQGTPLVTSS
E9, G7	PTIYFEDGRTSYYPGEAAFDI	WGQGTPLVTSS
F3, G4	PTIYFEDGRTSYYPGEAAFDI	WGQGTPLVTSS
A12, B5, B8, B9	TTAVTTTFDY	WGQGTPLVTSS
E8	GGHTVLSPLSNWFDP	WGQGTPLVTSS
F7	GYGLFDY	WGQGTPLVTSS
E11	DMGGAQRLPDH	WGQGTPLVTSS

Figure 5C

clones	SEQ ID NO.	FR1	CDR1	FR2
4D, 10C, 4G	219	GGGLVQPGASVKVCKASGY	TFSDYFMH	CVRQAPGQGLEWMGL
8A	220	RCPAKLLDT	PFSVYFMH	WVRQAPGQGLEWMGL
3G	221	RCPAKLLDT	PSGDIYFMH	WVRQAPGQGLEWMGL
1A	222	SGGLVQPGAKVLRISCVASGF	TFSSSAMS	WVRQAPGKGLEWVS
7H	223	LGS	PYSSSAMS	WVRQAPGKGLE?VSF
6F	224	VESGGVWPQGAKEVLRISCAASGF	SFEDYAMH	WVRQPPGKGLEWVAL
4F	225	AASGF	IFDDFAMH	WFQAVPGKGLQWVGL
5A	226	FWLGGPWRLSCAVSGY	TLSSSAMI	WVRQPPGKGLEFVS
1D	227	GGGLVQPGASLRISCVASGF	TLSSSAMS	CVRQAPGKGLEWVS
7E	228	WGRRGPANGVPVGSVPQPLGY	TFDDYAMH	WLRQIPGKGLQWVSL
9E	229	WTGGGVWPQGGSLRVSVAAAGY	TFDDYAMH	WLRQIPGKGLQWVSL
12B	230	AESGGGVWPQGGSLRLSCAASGF	TFSRYTIS	WVRQAPGKGLEWVS

Figure 6A

clones	CDR2	FR3
4D, 10C, 4G	VNPTNGYTAYAPKFQG	RVTMTQRQFTSTVYMELSSLRSEDTAVYFCAR
8A	VNPTNGYTAYAPKFQG	RVTMTQRQFTSTVYMELSSLRSEDTAVYFCAR
3G	VNPTNGYTAYAPKFQG	RVTMTQRQFTSTVYMELSSLRSEDTAVYFCAR
1A	ISGNGFSTYYADSVK	RFTISRDNKNTLYLQMNLSLRAEDTAEYYCTK
7H	IS?NGLSAYYADSVKG	RFTISRDNNS?NTVYLQMNLSLRSEDTAEYYCVK
6F	ISWDVISAYYADSVKG	RFTISRDNKNSLYLQMDSLRPEDSGLYYCGR
4F	MSWDGVSAYYADSVKG	RFTISRDNKKNALYLQMNLSLGVEDTALYFCAK
5A	ISGNGLSAYYADSVKG	RFTISRDNKNTVYLQMNLSLRAEDTAEYYCVK
1D	SSGNGFSAYYADSVKG	RFTISRDNKNTLYLQMNLSLVAEDTAEYYCTK
7E	LSWDGVSAYYADSVKG	RFTISRDNKNSLYLQMNLSLVAEDTALYFCAK
9E	LSWDGVSAYYADSVKG	RFTISRDN?KNSLYLQMNLSLIAEDTALYFCAK
12B	ISTDGGSTIYYTDSVKG	RFTISRDNKAKNSLSLQMNISLRDEDTAVYYCAR

Figure 6B

clones	CDR3	FR4
4D, 10C, 4G	VKSSDSIDAFDI	WGQGTMTVTSS
8A	VKSSDSIDAFDI	WGQGTMTVTSS
3G	VKSSDSIDAFDI	WGQGTMTVTSS
1A	VKYGSGSHFWFDP	WGQGTMTVTSS
7H	VYGSRSHF	
6F	DIGQRTMDV	WGQGTMTVTSS
4F	DMGGGLRFPHF	WGQGTMTVTSS
5A	VKYGSRSHFFDS	WGQGTMTVTSS
1D	VNYGSRSHFYFGS	WGQGTMTVTSS
7E	DMGGAQRLPDH	WGQGTMTVTSS
9E	DMGGAQRLPDH	WGQGTMTVTSS
12B	VFFGGNFRAHWFYDL	WGQGTMTVTSS

Figure 6C